Viral Infection Increases the Risk of Idiopathic Pulmonary Fibrosis: A

Meta-analysis

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Abstract

Background

Idiopathic pulmonary fibrosis (IPF) is a chronic, progressive, fibrotic lung disease with a poor prognosis. Although many factors have been identified to possibly trigger or aggravate IPF, such as viral infection, the exact etiology of IPF remains unclear. Until now, there has been no systematic review to quantitatively assess the role of viral infection in IPF.

Objective

This meta-analysis aims to present a collective view on the relationship between viral infection and IPF.

Methods

We searched studies reporting the effect of viral infection on IPF from Pubmed, Embase, Cochrane Library, Web of Science, and Wiley Online Library databases. The value of OR with 95% CI was calculated to assess the risk of virus in IPF. We also estimated statistical heterogeneity by I² and Cochran Q test, publication bias by funnel plot, Begg's test, Egger's test and trim and fill method. Regression, sensitivity and subgroup analyses were performed to assess the effects of confounding factors, such as gender and age.

Results

We analyzed 20 case-control studies with 1287 participants from 10 countries. The pooled OR of all viruses indicated that viral infection could significantly increase the risk of IPF (OR: 3.48, 95% CI: 1.61-7.52, p=0.001), but not that of

exacerbation of IPF (OR: 0.99, 95% CI: 0.46-2.12, p=0.988). In addition, all

analyzed viruses including Epstein-Barr virus (EBV), cytomegalovirus (CMV),

human herpesvirus 7 (HHV7) and human herpesvirus 8 (HHV8) were

associated with a significant elevation in the risk of IPF, except human

herpesvirus 6 (HHV6).

Conclusions

The presence of persistent or chronic, but not acute, viral infections including

EBV, CMV, HHV7 and HHV8, significantly increases the risk of developing IPF,

but not exacerbation of IPF. These findings imply that viral infection could be a

potential risk factor for IPF.

Key words: Idiopathic pulmonary fibrosis, viral infection, virus, meta-analysis.

Abbreviations

ATS, American Thoracic Society

CFA, cryptogenic fibrosing alveolitis

CMV, cytomegalovirus

CI, confidence intervals

EBV, Epstein-Barr virus

EMT, Epithelial Mesenchymal Transition

ERS, European Respiratory Society

HBoV, Human Boca virus subtype 1

HCV, hepatitis C virus

HHV6, human herpesvirus 6

HHV7, human herpesvirus 7

HHV8, human herpesvirus 8

ICTV, International Committee on Taxonomy of Viruses

IFN-α, interferon-α

IFN-γR-/-, IFN-γR-deficient mice

ILD, interstitial lung disease

IPF, idiopathic pulmonary fibrosis

KSHV, Kaposi's sarcoma-associated herpes virus

LMP1, latent membrane protein 1

MHV68, murine γ-herpesvirus 68

NOS, Newcastle-Ottawa Scale

OR, odds ratio

PCR, polymerase chain reaction

PRISMA, preferred reporting items for systematic reviews and meta-analysis

RSV, respiratory syncytial virus

TGFβ1, transforming growth factor-β1

TTV, torque teno virus

Introduction

Interstitial lung diseases (ILD) is a varying group of disorders with pulmonary parenchyma involvement; idiopathic pulmonary fibrosis (IPF) is one of the major idiopathic ILD. IPF, with an incidence of 2.8-9.3 per 100000 per year, is a chronic, progressive and fibrotic lung disease characterized by fibroblast proliferation, extracellular matrix accumulation and destruction of pulmonary architecture.^{1,2} It is prone to occur in men and those who are older than 50-years.³⁻⁵ In severe cases, it develops into restrictive pulmonary ventilatory dysfunction, impaired gas exchange, and even respiratory failure.⁶⁻⁸ The prognosis of IPF is poor, with the median survival after diagnosis generally estimated at 2-5 years, although it may be prolonged to 6.87-7.91 years under specific antifibrotic therapy.⁹⁻¹³ Although many studies have focused on IPF, the etiology of IPF still remains unclear.

In addition to genetic factors, ¹⁴⁻¹⁷ a variety of environmental exposures have been identified to be closely related to the initiation and progression of IPF, including cigarette smoking, metal and wood dusts exposure, silica and agricultural surroundings, and microbial infections. ¹⁸⁻²⁰ Among these factors, the relationship between virus and IPF is widely investigated, and accumulating evidence implies that viral infections may play an important role in the initiation and exacerbation of IPF. ^{21,22} However, the exact pathogenetic relationship between viral infection and IPF remains the subject of ongoing investigation. ^{1,4,23-25}

The purpose of this meta-analysis is to estimate the association between

viral infections and the development or exacerbation of IPF.

Methods

Search strategy and selection criteria

The search flow diagram of this meta-analysis is presented in **Figure 1**. This study was performed according to the standards set forth by the Preferred Reporting Items for Systematic Reviews and Meta-Analysis (PRISMA) statement.²⁶ In this meta-analysis, we searched Pubmed, Embase, Cochrane Library, Web of Science, Wiley Online Library databases, conference proceedings and other unpublished studies with terms "idiopathic pulmonary fibrosis", "IPF", "cryptogenic fibrosing alveolitis", "CFA", and "virus". All included studies were published in English up to December 31, 2018. In addition, the related articles in references were manually searched.

Inclusion criteria

We included controlled studies in English on patients diagnosed with IPF and acute exacerbation of IPF according to the American Thoracic Society (ATS) and European Respiratory Society (ERS) statements^{3,4,27,28} and the reports of Collard et al, respectively.^{29,30} Viral infection had to be detected by laboratory examinations, with the methodlogy of virus detection described in detailed. We only included studies with at least 3 participants in the control group.

Exclusion criteria

Uncontrolled studies or studies lacking data regarding the control group were

excluded. Studies were not eligible if all the participants were detected as virus positive/negative in both case and control groups, because the odds ratio (OR) could not be calculated.^{31,32} Studies that reported mean antibody titers instead of negative or positive results of virus detection were also eliminated. In addition, studies which recruited participants after lung or bone marrow transplantation were excluded.

After full-text assessments of 48 studies, a total of 28 were excluded. Six studies were excluded because they divided participants into case and control groups according to the presence or absence of virus infection instead of IPF diagnosis. Eight studies were excluded due to absence of control group. Four studies were excluded because virus detection outcomes were positive or negative in all participants. Another study were excluded because it only provided the levels of mean antibody titer. We further removed 9 studies where IPF diagnostic criteria did not refer to ATS/ERS statements. Finally, 20 eligible studies were included and analyzed in this meta-analysis.

Data extraction

Literature search and data extraction were independently performed by two researchers (Gaohong Sheng and Peng Chen). Disagreement between the two researchers were settled by arbitration of the principal investigator (Hui-Lan Zhang). The characteristics of studies and participants, and data of virus detection were collected and analyzed. Furthermore, the data of acute exacerbation of IPF were collected and analyzed as well.

Quality Assessment

The Newcastle-Ottawa Scale (NOS) was used to assess the quality of all included studies.³³ Each study was assessed by two researchers independently. The studies with a NOS score of 5 stars or more are considered as high quality and qualified to be included and analyzed in this meta-analysis. To estimate whether control groups were adequately selected, those ones that utilized healthy volunteers as control were rendered a star by NOS. For the item of comparability, studies which satisfied the most important factor would be given a star. One star would be appraised if the age gap between the case and control groups is less than or equal to 10 years. Since the second important factor is the sex ratio, one star would be given to a study where the value of sex ratio is within the range of 1/2 - 2 by control group dividing case group.

Statistical Analysis

We performed this meta-analysis using OR with 95% confidence intervals (CI) as the effect measure. In addition, in the studies with zero events in only one group, 0.5, the continuity correction was added to each cell of the table. Mantel–Haenszel method was used to estimate the pooled OR in order to reduce the bias. Methods of sample collection and viral detection varied among studies. We preferentially extracted results that used polymerase chain reaction (PCR) with lung tissue samples. The IgM level was used only when PCR data were not available.

We evaluated publication bias using funnel plot, Egger's regression asymmetry test and Begg's adjusted rank correlation test. When obvious publication bias was found, we used trim and fill method to adjust and confirm our results. The heterogeneity of study was assessed using Cochran'Q statistic, and its magnitude was estimated using I^2 statistic. Random effect model was used when significant heterogeneity was observed (p < 0.1 or I^2 > 50%), otherwise we used fixed effect model. Sensitivity analyses was performed to examine stability of pooled outcome. In addition, Meta regression and subgroup analyses were performed according to the mean age of patients and the proportion of man.

All comparisons were two-tailed, and p < 0.05 was recognized as statistical significance. STATA software 12.0 was used to perform statistical analyses.

Results

Characteristics of studies and patients

This meta-analysis reviewed and analyzed 20 case-control studies published from 1984 to 2018, covering 1287 participants from 10 countries (**Table 1**) (634 IPF cases and 653 controls). The mean age of participants was 59.75 years old. The mean proportion of males was 69% (**Table 2**). More detailed characteristics of the studies and patients are shown in **Table 1** and **Table 2**. All included studies were distinguished as high-quality studies by NOS quality assessments (**Supplementary Table 1**). According to the virus classification of International Committee on Taxonomy of Viruses (ICTV), a total of 19 virus species were detected using various laboratory examinations. Among them, five species of virus had more than two sets of data, including Epstein-Barr virus (EBV), cytomegalovirus (CMV), human herpesvirus 6 (HHV6), human herpesvirus 7 (HHV7), and human herpesvirus 8 (HHV8) (**Supplementary Table 2**).

Relationship between virus infection and IPF

All viruses

For all species of virus, we analyzed the data of viral infection in 634 patients from 20 studies. A significant elevation associated with viral infection was found in the risk of IPF (OR: 3.48, 95% CI: 1.61-7.52, p=0.001) (**Figure 2a**). The heterogeneity was I²: 76.0% (p<0.001). Publication bias was found in the funnel plot (**Supplementary Figure 1a**). The p value of Begg's test was 0.004

(Supplementary Figure 1b), while that of Egger's test p=0.001 (Supplementary Figure 1c). Considering the influence of publication bias, we conducted trim and fill method (Supplementary Figure 1d) and the result of OR (1.64, 95% CI: 0.73-3.69, p=0.231) failed to confirm the stability of this effect.³⁶ Almost all studies published before 2000 were removed due to inability to apply ATS/ERS guidelines, which likely led to this publication bias. Sensitivity analysis examined the robustness by evaluating the impacts of small sample studies (Supplementary Figure 1e) and individual studies (Supplementary Figure 1f) on pooled effects.

EBV

EBV is a typical species of the Gammaherpesvirinae family. A total of 10 studies consisting of 252 patients reported data on EBV infection, and EBV infection was associated with a significant elevation in the risk of IPF (OR: 9.83, 95% CI: 4.22-22.91, p<0.001) (Figure 2b). The degree of the heterogeneity was I²: 41.5% (p=0.081). Publication bias was not negligible from the funnel plot (**Supplementary Figure 2a**), and also detected by Begg's test (p=0.032) **2b**) as well as (Supplementary Figure Egger's test (p=0.030)(Supplementary Figure 2c). Considering the influence of publication bias, we conducted trim and fill method (Supplementary Figure 2d) and the result of OR (6.16, 95% CI: 2.74-13.86, p<0.001) further confirmed the stability of our results. No obvious influences were found from small sample studies (Supplementary Figure 2e) or individual studies through sensitivity analyses

(Supplementary Figure 2f).

CMV

CMV is a common virus which belongs to the Betaherpesvirinae family, and a total of 209 patients from 9 studies were analyzed. We evaluated the pooled effect and found infection of CMV contributing significantly to IPF risk (OR: 2.45, 95%CI: 1.29-4.66, p=0.006) (Figure 2c). The degree of the heterogeneity was I²: 24.1% (p=0.229). No obvious publication bias was observed from funnel plot (Supplementary Figure 3a), quantitative Begg's test (p=0.754) (Supplementary Figure 3b) or Egger's test (p=0.355) (Supplementary Figure 3c). Neither small sample studies (Supplementary Figure 3d) nor individual studies (Supplementary Figure 3e) had significant impacts on the pooled OR by sensitivity analysis.

HHV6

Three studies involving 80 patients provided data on HHV6, which also belongs to the Betaherpesvirinae family. The result of pooled OR showed no statistically significant elevation in risk of IPF (OR: 3.16, 95%CI: 0.41-24.50, p=0.270) (**Figure 2d**). The degree of the heterogeneity was I²: 71.4% (p=0.030). The funnel plot (**Supplementary Figure 4a**) showed no evident publication bias by qualitative evaluation, and the quantitative assessments via Begg's test (p=1.000) (**Supplementary Figure 4b**) and Egger's test (p=0.242) (**Supplementary Figure 4c**). Sensitivity analysis did not show prominent influences on pooled effect from small sample studies (**Supplementary**

Figure 4d) or individual studies (Supplementary Figure 4e).

HHV7

Another member of Betaherpesvirinae family, HHV7 was investigated in 4 studies with 95 patients. Pooled effect showed patients with HHV7 infection had higher risk of IPF (OR: 2.48, 95%CI: 1.14-5.40, p=0.022) (Figure 2e). The degree of the heterogeneity was I²: 0.0% (p=0.625). No publication bias was found by funnel plot (Supplementary Figure 5a), Begg's test (p=0.308) (Supplementary Figure 5b) and Egger's test (p=0.702) (Supplementary Figure 5c). Sensitivity analysis showed no significant influences on overall effect, including small sample studies (Supplementary Figure 5d) and individual studies (Supplementary Figure 5e).

HHV8

HHV8 is a species which belongs to the Gammaherpesvirinae family, also known as Kaposi's sarcoma-associated herpes virus (KSHV).³⁷ We found 2 studies of HHV8 and 1 study of KSHV to obtain the overall effect of OR, and the pooled effect showed that HHV8 infection increased the risk of IPF significantly (OR: 8.88, 95%CI: 2.70-29.14, p<0.001) (Figure 2f). The degree of heterogeneity was I²: 0.0% (p=0.662). No publication bias was found by the funnel plot (Supplementary Figure 6a), Begg's test (p=1.000) (Supplementary Figure 6b) or Egger's test (p=0.508) (Supplementary Figure 6c). Sensitivity analysis for overall effect showed no significant influences from small sample studies (Supplementary Figure 6d) or

individual studies (Supplementary Figure 6e).

Relationship between multiple viral infections and IPF

Six studies containing 174 patients provided data on multiple viral infections. The result of pooled OR did not show that multiple viral infections increased risk of IPF significantly (OR: 3.37, 95%CI: 0.69-16.44, p=0.133) (Figure 3a). The degree of the heterogeneity was I²: 60.4% (p=0.027). The funnel plot (Supplementary Figure 7a) showed no apparent publication bias by qualitative evaluation and the quantitative assessments by Begg's test (p=0.707) (Supplementary Figure 7b) and Egger's test (p=0.284) (Supplementary Figure 7c). Sensitivity analysis showed that influences on pooled effect were not remarkable from small sample studies (Supplementary Figure 7d) or individuals studies (Supplementary Figure 7e).

Relationship between viral infection and acute exacerbation of IPF

A total of 3 studies provided data on acute exacerbation of IPF patients. However, viral infection was not found to associate with acute exacerbation of IPF (OR: 0.99, 95%CI: 0.46-2.12, p=0.988) (Figure 3b). The degree of heterogeneity was I²: 0.0% (p=0.759). No publication bias was found in the funnel plot (Supplementary Figure 8a), Begg's test (p=0.296) (Supplementary Figure 8b) and Egger's test (p=0.209) (Supplementary Figure 8c). Sensitivity analysis showed no obvious influences caused by small sample studies (Supplementary Figure 8d) or individual studies (Supplementary Figure 8e).

Meta regression

Age and gender were used as covariates for Meta regression analyses and the results suggested that the viral infection related to IPF in an age and gender independent manner (p>0.05 for all) (Supplementary Table 3).

Subgroup analyses

We also performed subgroup analyses based on the mean age of patients and the proportion of males for all viruses, EBV CMV and viruses of multiple infections, with no significant difference between subgroups was found (Supplementary Table 4).

Discussion

This meta-analysis including 20 case-control studies with 634 IPF cases and 653 controls, covering 19 virus species, aimed to evaluate the relationship between viral infection and IPF. We found that viral infection was associated with higher risk of IPF but not of acute exacerbation of IPF. Further, we performed subgroup analyses to evaluate the relationship between IPF and each kind of virus, including EBV, CMV, HHV6, HHV7 and HHV8. The results showed that EBV, CMV, HHV7 and HHV8, but not HHV6, were associated with a significant increase in the risk of IPF.

So far, the exact etiology and pathogenesis of IPF still remains unclear. Accumulating studies have focused on the role of viral infection in IPF patients, but the interpretation of them was hindered by their relatively small sample size and differing populations studied.^{21,22,38,39}

Historically, "IPF" comprises a heterogeneous group of different histological and clinical entities arising in an idiopathic setting. Before ATS/ERS refined IPF as those patients with usual interstitial pneumonia (UIP) pattern on lung biopsy or high-resolution computed tomography (HRCT), the diagnosis of IPF was imprecise and variable, 3,4,6,27 which is a major pitfall of this meta-analysis. Therefore, we compared the outcomes between all studies related to "IPF" and those referring to ATS/ERS statements in **Supplementary Table 5**. There were no dramatic changes in any results for all viruses, EBV, CMV, HHV6, HHV7 and HHV8. In addition, the stability of the result of all studies involving "IPF" for all viruses was confirmed by the trim and fill method, while it was not

confirmed in the result of studies using the ATS/ERS criteria. This implies that the publication bias of studies using the ATS/ERS criteria might influence the overall results. The "IPF" patients in those excluded studies might accord with ATS/ERS statements, since they were all published earlier than publication of these statements. The characteristics of studies involving "IPF" but not definitely referring to ATS/ERS statements are shown in **Supplementary Table 6**.

We further analyzed and calculated OR for patients with multiple chronic viral infections but did not find a statistically significant increase in the risk of IPF. The species and amount of viruses detected in different studies could result in bias, as humans are likely to harbor multiple persistent viruses without obvious symptoms throughout their lifetimes.⁴⁰ Thus, further study are needed to detect more viruses in IPF patients in order to assess the effect of multiple viral infections on IPF.

The result of subgroup analysis of age suggested the possibility that younger participants with viral infection might exhibit higher risk of IPF, although the interaction term for age (p=0.186) was not significant. It seemed that IPF of the young might be influenced by factors that differ from the elderly. The changes in the lung with age could facilitate the development of lung disease, including pulmonary fibrosis and lung infections.⁴¹ Moreover, Naik et al suggested that aged lung may be particularly susceptible to viral-induced fibrosis.³⁸ However, the role of age is still poorly understood in viral infection

related IPF, and more works are needed to clarify in the future.

Previous studies have reported that murine models of viral infection could foster lung fibrogenesis. However, the murine models predominantly focus on the role of acute inflammation driven by viral infection. Virgin et al have compared the differences between chronic and acute viral infections. Thus, it should be noted that the majority of the viruses analyzed in our study are chronic/persistent viruses that are likely to have been acquired prior to the development of IPF. In view of this, we must be cautious regarding the differences in the pathogenesis of pulmonary fibrosis between humans and murine models. More appropriate animal models are requisite to further reveal the pathogenetic relationship between viral infection and IPF.

Malizia et al suggested that EBV-infected epithelial cell may play a role in IPF via transforming growth factor- β 1 (TGF β 1) and CUX1/Wnt signaling pathway, which were involved in Epithelial Mesenchymal Transition (EMT), the mechanism of fibrosis production in tissue. Human Boca virus subtype 1 (HBoV) might trigger development of pulmonary fibrosis by regulating the expression of cytokines. It was also reported that latent membrane protein 1 (LMP1) expressed by EBV could induce EMT in synergy with TGF β 1. Meanwhile, our results showed that infection of EBV was associated with a significant elevation in the risk of IPF, in agreement with previous studies.

Studies have reported that antiviral drugs against EBV and herpesvirus could stabilize the condition of patients with IPF. 16,47 Additionally, interferon- α

(IFN- α) treatment revealed substantial anti-fibrotic effects in a mouse model.⁴⁸ These studies hint that specific therapy towards culpable viral infections might be of therapeutic use.

All studies included in this meta-analysis are retrospective case-control studies with limitation of explaining causal relationships. The sample size in some subgroups is small. In addition, the viral infection rates are variable due to the sites of biological samples and multiple virus detection techniques. Different methods lead to alterations in sensitivity and specificity. Given these challenges, larger-scale samples and higher quality studies are needed in the future to draw conclusions on the exact causal relationship between the virus and IPF and/or virus and acute exacerbation of IPF.

In conclusion, the presence of viral infections associated with a significant increase in the risk of IPF, specifically for EBV, CMV, HHV7 and HHV8 infections. Our study supports the hypothesis that virus infection could play a key role in pathogenesis of IPF.

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Competing interests

The authors declare that they have no competing interests.

Authors' contributions

Hui-Lan Zhang was the guarantor of the entire manuscript for designing and supervising the entire study. Gaohong Sheng and Peng Chen independently completed the literature retrieval and data extraction. The manuscript was drafted by Gaohong Sheng and checked and approved by Wanguang Zhang. Yanqiu Wei addressed grammatical/wording issues of the manuscript to smooth it out. Huihui Yue contributed the tables and figures. Jiaojiao Chu was in charge of statistical methods in this study. Jianping Zhao and Yihua Wang provided guidance for writing the part of Discussion.

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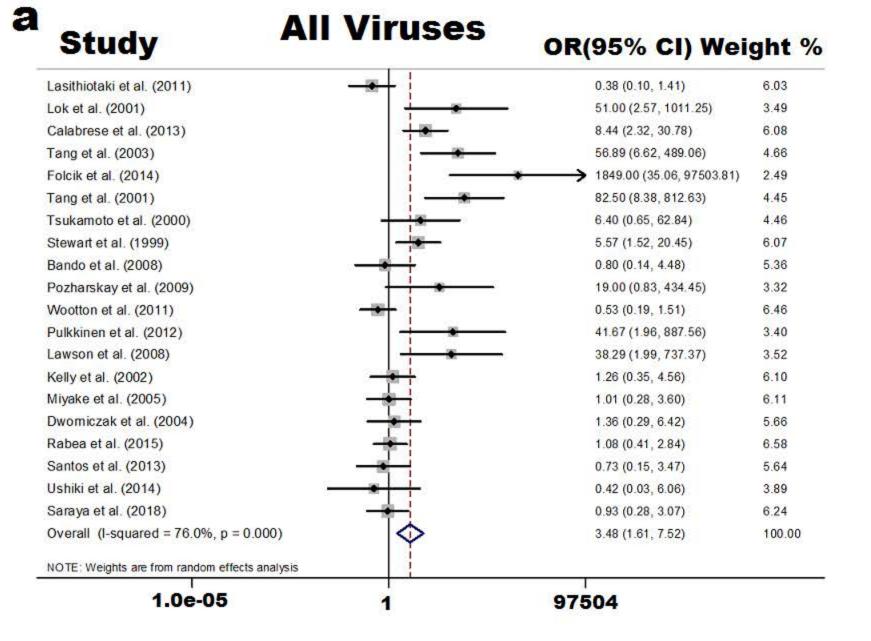
Figure Legends

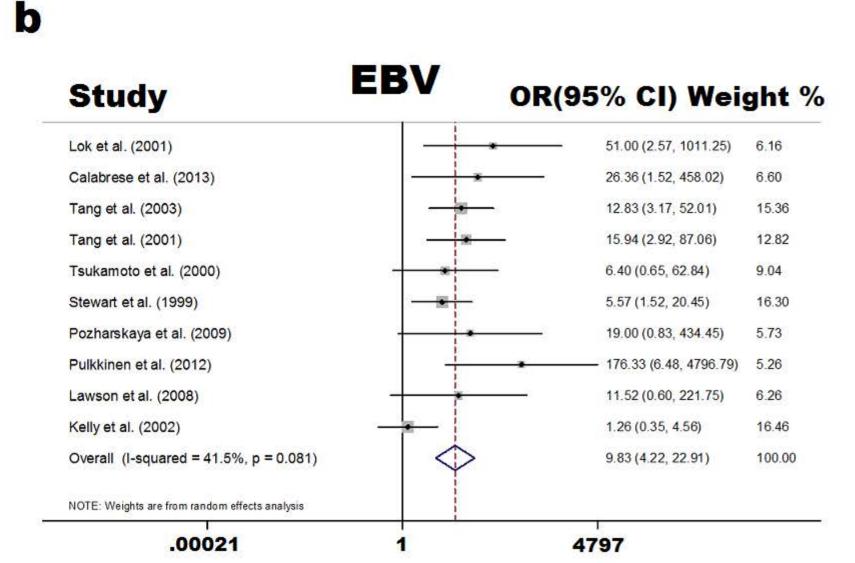
Figure 1. Search flow diagram for included studies in the meta-analysis.

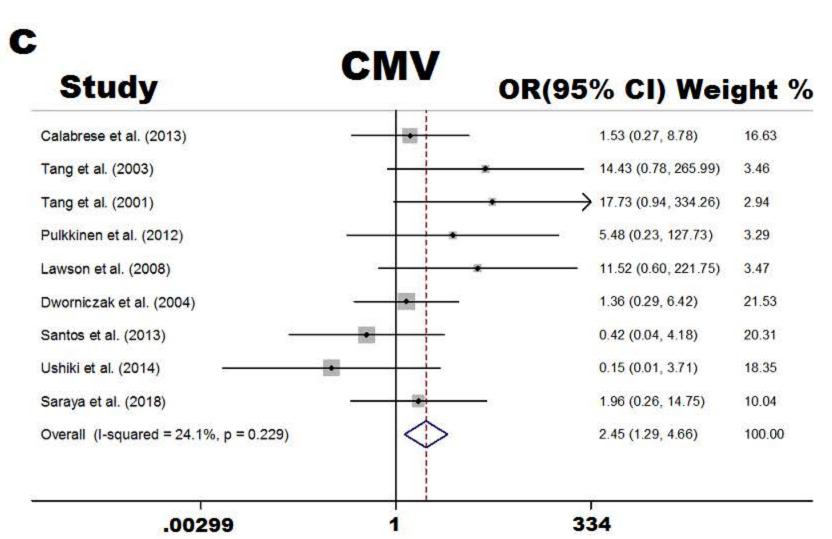
Figure 2. Forest plot shows the pooled effect of OR for viral infection in idiopathic pulmonary fibrosis (IPF) patients and non-IPF controls. **a.** Viral infection was associated with a significant increase in the risk of IPF (OR: 3.48, 95%CI: 1.61-7.52, p=0.001). **b.** EBV infection was associated with significantly higher risk of IPF (OR: 9.83, 95%CI: 4.22-22.91, p<0.001). **c.** CMV infection significantly increased the risk of IPF (OR: 2.45, 95%CI: 1.29-4.66, p=0.006). **d.** No significant difference was found in HHV6 infection between IPF patients and controls (OR: 3.16, 95%CI: 0.41-24.50, p=0.270). **e.** HHV7 infection showed significantly higher risk of IPF (OR: 2.48, 95%CI: 1.14-5.40, p=0.022). **f.** HHV8 infection increased the risk of IPF (OR 8.88, 95%CI: 2.70-29.14, p<0.001).

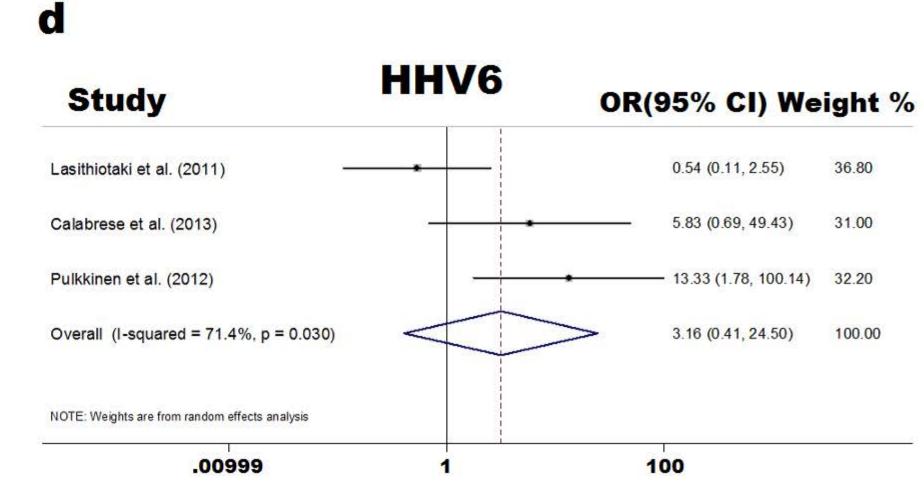
Figure 3. Forest plot shows the pooled effect of OR for viral infection in idiopathic pulmonary fibrosis (IPF) patients with acute exacerbation and with multiple viral infections compared to non-IPF controls. **a.** The pooled effect showed that viral infection was not associated with acute exacerbation of IPF (OR: 0.99, 95%CI: 0.46-2.12, p=0.988). **b.** No significant increase was found in multiple viral infections between IPF patients and controls (OR: 3.16, 95%CI: 0.41-24.50, p=0.270).

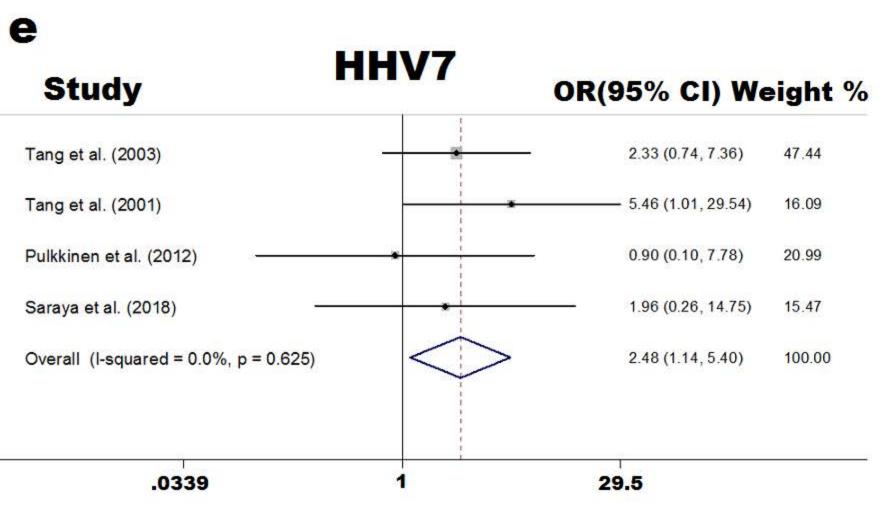
and the same of th		
Search strategy (idiopathic pulmonary fibrosis OR IPF OR cryptogenic fibrosing alveolitis OR CFA) AND (virus)	Database Pubmed Embase Cochrane library Web of science	372 633 1092 1892
	Wiley online library	385
Duplica	ates removed 1536	
Title and abstract screened 2838		
	Articles excluded	2790
	Not in English	325
	Not in humans	833
	IPF or CFA means other abbreviations	479
	Reviews or editorials	237
	Not related to target topic	916
Full-text assessment 48	-	
	Articles excluded	19
	IPF not as case group	6
	Only IPF but no control	8
	Only negative or positive results	4
	Only mean titer of antibody	1
Diagnosis criteria assessment 29		
	Not refer to ATS/ERS guidelines	9
Articles included 20		

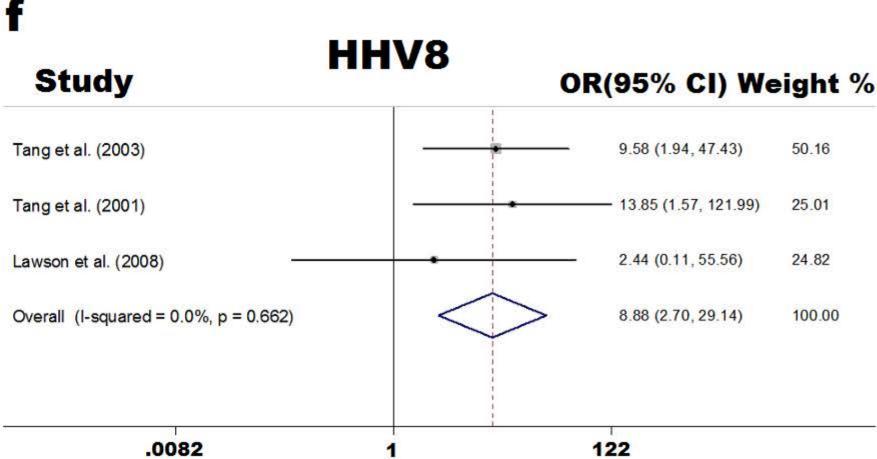












All Viruses Study **All Viruses** OR(95% CI) Weight % Study OR(95% CI) Weight % 0.22 (0.01, 5.85) 12.92 Lasithiotaki et al. (2011) 45.04 1.51 (0.13, 17.24) 17.04 Wootton et al. (2011) 1.22 (0.41, 3.58) Calabrese et al. (2013) Tang et al. (2003) 15.61 (3.15, 77.41) 22.08 Ushiki et al. (2014) 12.81 0.42 (0.03, 6.06) 50.00 (3.88, 643.87) 16.37 Pulkkinen et al. (2012) Saraya et al. (2018) 0.93 (0.28, 3.07) 42.15 3.59 (0.17, 76.09) 13.84 Lawson et al. (2008) 100.00 Overall (I-squared = 0.0%, p = 0.759) 0.99 (0.47, 2.12) 17.74 Saraya et al. (2018) 0.62 (0.06, 6.22) Overall (I-squared = 60.4%, p = 0.027) 3.37 (0.69, 16.44) 100.00 NOTE: Weights are from random effects analysis 34.9 .0286 .00155 644

Table 1. Characteristics of included studies.

	Study	Assessment			
Study	design	NOS	Diagnosis	Country	Year
Lasithiotaki et al. 2011	Case-control	7	ATS/ERS statement, 2000	Greece	NA
Lok et al. 2001	Case-control	7	UIP pattern on histopathology and HRCT	UK	NA
Calabrese et al. 2013	Case-control	8	ATS/ERS statement, 2002	Italy	1998-2010
Tang et al. 2003	Case-control	8	UIP pattern on histopathology	USA	NA
Folcik et al. 2014	Case-control	7	ATS/ERS/JRS/ALAT statement, 2011	USA	NA
Tang et al. 2001	Case-control	6	UIP pattern on histopathology	USA	NA
Tsukamoto et al. 2000	Case-control	7	UIP pattern on histopathology	Japan	NA
Stewart et al. 1999	Case-control	8	UIP pattern on histopathology and HRCT	UK	NA
Bando et al. 2008	Case-control	6	ATS/ERS statement, 2000	Japan	1991-2003
Pozharskaya et al. 2009	Case-control	7	UIP pattern on histopathology	USA	NA

Wootton et al. 2011	Case-control	6	ATS/ERS statement, 2000 Collard et al. 2007	Korea; Japan	NA
Pulkkinen et al. 2012	Case-control	7	ATS/ERS statement, 2000/2002	Finland	NA
Lawson et al. 2008	Case-control	7	ATS/ERS statement, 2000/2002	USA	NA
Kelly et al. 2002	Case-control	7	UIP pattern on histopathology and HRCT	UK	NA
Miyake et al. 2005	Case-control	7	ATS/ERS statement, 2002	Japan	2001
Dworniczak et al. 2004	Case-control	6	ATS/ERS statement, 2000	Poland	NA
Rabea et al. 2015	Case-control	7	ATS/ERS statement, 2002	Egypt	NA
Santos et al. 2013	Case-control	7	ATS/ERS statement, 2002	Brazil	1993-2011
Ushiki et al. 2014	Case-control	7	ATS/ERS statement, 2002 Collard et al. 2007	Japan	2007-2011
Saraya et al. 2018	Case-control	8	Collard et al. 2007/2016	Japan	2012-2015

NOS, Newcastle-Ottawa Scale; ATS, American Thoracic Society; ERS, European Respiratory Society; JRS, Japanese Respiratory Society; ALAT, Latin American Thoracic Society; UIP, usual interstitial pneumonitis; UK, United Kingdom; USA, the United States of America; NA, not available; HRCT, high resolution computed tomography

Table 2. Characteristics of included participants.

									Immuno-
				Patients	Controls	Age,	Gender,	Smoking/	suppression/
Study	Virus	Sample	Method	positive/total	positive/total	у	M/F	total	total
1 141-1-4-1-1 -4 -1	HSV-1	lung	PCR	1/11	0/4	NA	NA	NA	NA
Lasithiotaki et al.	HHV-6	BALF	PCR	5/13	8/13	04.05	40/0	4.4/00	NIA
2011	CMV/HHV7/8	lung/BALF	PCR	0/24	0/17	64.95	18/8	14/26	NA
Lok et al. 2001	EBV	lung	PCR and IHC	8/14	0/19	55.55	19/14	NA	NA
	EBV			13/55	0/41				
Calabrese et al.	HHV-6	I		7/55	1/41	52	47/30	53/77	NIA
2013	CMV	lung	PCR and IHC	4/55	2/41				NA
	PVB19			0/55	1/41				
	EBV			21/33	3/25				
Tang et al.	CMV	lung	PCR	7/33	0/25	52.41	32/18	NA	30/45
2003	HHV-7	lung	PCR	14/33	6/25	52.41	32/10		30/45
	HHV-8			15/33	2/25				
Folcik et al. 2014	HVS	lung	ISH and IHC	21/21	0/21	61.3	29/13	NA	NA
	CMV			7/23	0/19				
	EBV			15/23	2/19				
Tang et al.	HHV-7			9/23	2/19	NA	NA	NA	
2001	HHV-8	lung	PCR	10/23	1/19				NA
	HSV-1/2 VZV/HHV-6			0/23	0/19				

Tsukamoto et al. 2000	EBV	lung	PCR	24/25	15/19	57.33	25/9	21/29	0/29
Stewart et al. 1999	EBV	lung	PCR	13/27	4/28	55.47	37/18	37/55	16/27
Bando et al. 2008	TTV	serum	PCR	55/57	138/142	65.29	147/52	153/199	0/199
Pozharskaya et al. 2009	EBV	lung	PCR	9/13	0/4	NA	NA	NA	NA
Wootton et al. 2011	TTV	BAL	PCR	12/83	7/29	64.06	87/25	80/112	NA
	EBV			11/12	0/11				
	HHV-6B	lung	PCR	10/12	3/11	58.71	18/3		
Pulkkinen et al.	HHV-7			2/12	2/11			11/21	2/21
2012	CMV			2/12	0/11			11/21	2/2 1
	HSV-1			1/12	0/11				
	HSV-2			1/12	0/11				
Lawson et al.	EBV			8/23	0/10				
2008	CMV	lung	IHC	8/23	0/10	NA	NA	NA	NA
2000	KSHV			2/23	0/10				
Kelly et al. 2002	EBV	serum	PCR	23/27	41/50	52.25	15/11	NA	36/77
Miyake et al. 2005	HCV	medical history	NA	7/104	4/60	NA	149/15	129/164	NA
Dworniczak et al. 2004	CMV	serum; blood leukocytes; BAL cells	PCR	12/16	11/16	38.85	21/11	14/32	0/32

Rabea et al.	HCV	serum	ELISA	9/30	17/60	50.55	38/52	19/90	NA				
2015													
Santos et al.	MeV	lung	IHC	2/13	3/24	61.16	18/19	NA	NA				
2013	CMV	lulig	1110	1/13	4/24	01.10	10/19	IVA	INA				
Ushiki et al.	RSV-B	DAI	PCR	1/7	0/7	60 F	11/3	NA	1/14				
2014	CMV	BAL	PCR	0/7	2/7	69.5		INA	1/14				
	HHV-7			3/27	5/51								
	CMV			2/27	2/51								
	Influenza			0/07	2/54								
Saraya et al.	AH3	sputum; BALF;		DOD	DOD	DOD	DOD	DOD	0/27	3/51	74.05	40/00	40/70
2018	Influenza	nasal swab	PCR	0/07	4/54	74.65	46/32	42/78	73/78				
	H1N1			0/27	1/51								
	HPIV			1/27	1/51								
	HMPV			0/27	1/51								

y, year; M/F, male/female; NA, not available; PCR, polymerase chain reaction; BALF, bronchoalveolar lavage fluid; IHC, immunohistochemistry; ISH, in situ hybridization; BAL, bronchoalveolar lavage; ELISA, enzyme linked immunosorbent assay; HSV, herpes simplex virus; HHV, human herpesvirus; CMV, cytomegalovirus; EBV, Epstein-Barr virus; PV, parvovirus; HVS, herpesvirus saimiri; VZV, varicella zoster virus; TTV, torque teno virus; KSHV, Kaposi's sarcoma herpesvirus; HCV, hepatitis C virus; MeV, measles virus; RSV, respiratory syncytial virus; AH3, influenza virus type A Hemagglutinin 3; H1N1, Hemagglutinin 1 Neuraminidase 1; HPIV, human parainfluenza virus; HMPV, human metapneumovirus

Supplementary Table 1. Newcastle-Ottawa Scale of quality assessment for included studies.

Study	Selection				Comparability [#]		Exposure	Star	
		Comparability of							
					cases and		Same method of		
	Is the case		Selection		controls on the		ascertainment		
	definition	Representativeness	of	Definition of	basis of the design	Ascertainment	for cases and	Non-Respon	
	adequate?	of the cases	Controls	Controls	or analysis	of exposure	controls	se rate	
Lasithiotaki et al. ¹ 2011	*	*	-	*	-	*	*	*	6
LOK et al. ² 2001	*	*	-	*	*	*	*	*	7
Calabrese et al. ³ 2013	*	*	-	*	*	*	*	*	7
Tang et al. ⁴ 2003	*	*	-	*	*	*	*	*	7
Folcik et al. ⁵ 2014	*	*	-	*	**	*	*	*	8
Tang et al. ⁶ 2001	*	*		*	-	*	*	*	6
Tsukamoto et al. ⁷	*	*	-	*	*	*	*	*	7
Stewart et al. ⁸ 1999	*	*	-	*	**	*	*	*	8
Bando et al. ⁹	*	*	-	*	**	*	*	*	8

Pozharskaya et al. ¹⁰ 2009	*	*	-	*	-	*	*	*	6
Wootton et al. ¹¹ 2011	*	*	-	*	**	*	*	*	8
Pulkkinen et al. ¹² 2012	*	*	-	*	*	*	*	*	7
Lawson et al. ¹³ 2008	*	*	-	*	-	*	*	*	6
Kelly et al. ¹⁴ 2002	*	*	-	*	*	*	*	*	7
Miyake et al. ¹⁵ 2005	*	*	-	*	*	*	*	*	7
Dworniczak et al. ¹⁶ 2004	*	*	-	*	*	*	*	*	7
Rabea et al. ¹⁷ 2015	*	*	-	*	**	*	*	*	8
Santos et al. ¹⁸ 2013	*	*	-	*	**	*	*	*	8
Ushiki et al. ¹⁹ 2014	*	*	-	*	*	*	*	*	7
Saraya et al. ²⁰ 2018	*	*	-	*	**	*	*	*	8

[#] One star will be given if the age gap is less than or equal to 10 years between the case and control groups, and another star means the sex ratio (male/female) in case and control groups within 1/2-2; IPF, idiopathic pulmonary fibrosis

Supplementary Table 2. Classification of viruses from included studies*.

Virus	N	Abbreviation	Order	(Sub)family	Genus	Species
Epstein-Barr virus	10	EBV	Herpesvirales	Gammaherpesvirinae	Lymphocryptovirus	Human gammaherpesvirus 4
Cytomegalovirus	10	CMV	Herpesvirales	Betaherpesvirinae	Cytomegalovirus	Human betaherpesvirus 5
Human herpesvirus 7	5	HHV7	Herpesvirales	Betaherpesvirinae	Roseolovirus	Human betaherpesvirus 7
Human herpesvirus 6	4	HHV6	Herpesvirales	Betaherpesvirinae	Roseolovirus	Human herpesvirus 6
Human herpesvirus 8	3	HHV8	Herpesvirales	Gammaherpesvirinae	Rhadinovirus	Human gammaherpesvirus 8
Kaposi's sarcoma herpes virus	1	KSHV	Herpesvirales	Gammaherpesvirinae	Rhadinovirus	Human gammaherpesvirus 8
Herpes simplex virus type 1	3	HSV1	Herpesvirales	Alphaherpesvirinae	Simplexvirus	Human alphaherpesvirus 1
Herpes simplex virus type 2	2	HSV2	Herpesvirales	Alphaherpesvirinae	Simplexvirus	Human herpesvirus 2
Hepatitis C virus	2	HCV	Unassigned	Flaviviridae	Hepacivirus	Hepacivirus C
Torque teno virus	2	TTV	Unassigned	Anelloviridae	Alphatorquevirus	Torque teno virus
Herpesvirus saimiri	1	HVS	Herpesvirales	Gammaherpesvirinae	Rhadinovirus	Saimiriine herpesvirus 2
Measles virus	1	MeV	Mononegavirales	Paramyxoviridae	Morbillivirus	Measles morbillivirus
Respiratory syncytial virus	1	RSV	Mononegavirales	Pneumoviridae	Orthopneumovirus	Human orthopneumovirus
Human parvovirus B19	1	PVB19	Parvoviridae	Parvovirinae	Erythroparvovirus	Primate erythroparvovirus 1
Varicella zoster virus	1	VZV	Herpesvirales	Alphaherpesvirinae	Varicellovirus	Human alphaherpesvirus 3

Influenza virus AH3	1	-	Unassigned	Orthomyxoviridae	Alphainfluenzavirus	Influenza A virus
Influenza virus H1N1	1	-	Unassigned	Orthomyxoviridae	Alphainfluenzavirus	Influenza A virus
Human parainfluenza virus	1	HPIV	Mononegavirales	Paramyxoviridae	Respirovirus; or Rubulavirus	Human rubulavirus 1/2/3/4
Human metapneumovirus	1	HMPV	Mononegavirales	Pneumoviridae	Metapneumovirus	Human metapneumovirus

^{*} Virus classification according to the report of International Committee on Taxonomy of Viruses (ICTV); N, the number of studies involved with this specific virus

Supplementary Table 3. Meta regression analyses of viral infection in IPF for age and gender.

P value

Virus	Age	Gender	Age and Gender
All viruses	0.328	0.959	0.484
EBV	0.312	0.239	0.726
CMV	0.741	0.663	0.936
HHV6	0.483	0.715	NA
HHV7	0.889	0.611	NA
HHV8	NA	NA	NA

IPF, idiopathic pulmonary fibrosis; EBV, Epstein-Barr virus; CMV, cytomegalovirus; HHV, human herpesvirus; NA, not available due to the insufficient number of studies

Supplementary Table 4. Subgroup analyses of OR for viral infection in participants with or without IPF.

	Studies/participants,			P value for
	n/N	OR [95% CI]	P value	interaction
		All viruses		
Age, y				
<60	9/508	5.24[2.00-13.75]	0.001	0.168
≥60	7/523	0.99[0.31-3.13]	0.986	0.100
Gender, M/N				
<0.65	7/469	3.09[1.00-9.52]	0.050	0.697
≥0.65	10/726	2.09[0.72-6.09]	0.177	0.097
		EBV		
Age, y				
<55	3/231	6.13[0.90-41.84]	0.064	0.050
≥55	4/155	14.49[3.26-64.35]	<0.001	0.656
Gender, M/N				
<0.65	4/264	9.06[1.55-52.86]	0.014	0.010
≥0.65	3/122	11.33[2.06-62.36]	0.005	0.910
		CMV		
Age, y				
<60	4/209	2.73[1.06-7.06]	0.038	0.400
≥60	3/129	0.63[0.18-2.27]	0.483	0.196
Gender, M/N				
<0.65	4/269	2.05[0.81-5.18]	0.129	0.524
≥0.65	3/69	1.16[0.37-3.62]	0.799	0.521
	M	ultiple viral infectio	ns*	
Age, y				
<60	3/177	10.86[1.82-64.71]	0.009	0.540
≥60	2/119	0.44[0.07-2.90]	0.393	0.516

Gender, M/N

<0.65	3/232	2.05[0.02-758.61]	0.327	0.996
≥0.65	2/64	2.83[0.35-22.78]	0.631	0.990

OR, odds ratio; IPF, idiopathic pulmonary fibrosis; y, years; M/N, men/total; EBV, Epstein-Barr virus; CMV, Cytomegalovirus; * two or more viruses simultaneously detected as positive in one participant

Supplementary Table 5. Comparison of results between studies involving "IPF" and those referring to ATS/ERS statements.

OR [95% CI], P value

Virus	Studies involving "IPF"	Studies referring to ATS/ERS guidelines
All viruses	3.01 [1.59-5.68], p=0.001	3.48 [1.61-7.52], p=0.001
EBV	4.93 [1.89-12.87], p=0.001	9.83 [4.22-22.91], p<0.001
CMV	2.45 [1.29-4.66], p=0.006	2.45 [1.29-4.66], p=0.006
HHV6	3.16 [0.41-24.50], p=0.270	3.16 [0.41-24.50], p=0.270
HHV7	2.48 [1.14-5.40], p=0.022	2.48 [1.14-5.40], p=0.022
HHV8	8.88 [2.70-29.14], p<0.001	8.88 [2.70-29.14], p<0.001

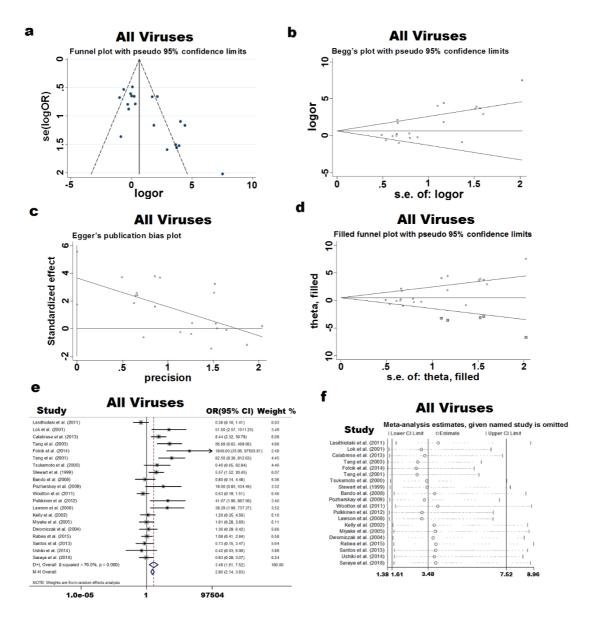
OR, odds ratio; IPF, idiopathic pulmonary fibrosis; ATS, American Thoracic Society; ERS, European Respiratory Society; EBV, Epstein-Barr virus; CMV, cytomegalovirus; HHV, human herpesvirus

Supplementary Table 6. Characteristics of studies and participants not referring to ATS/ERS statements.

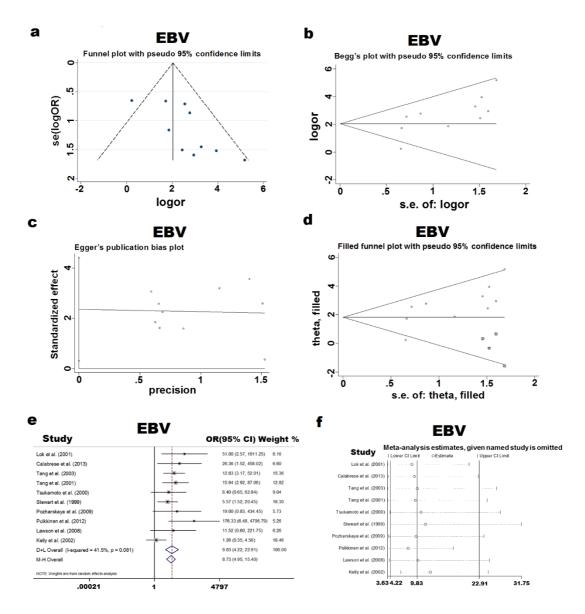
		Assessment				Patients	Controls	Age,	Gender,
Study	Diagnosis	NOS	Virus	Sample	Method	positive/total	positive/total	у	M/F
Wangoo et al. ²¹ 1997	Histology	6	EBV	lung	IHC, ISH, and PCR	0/12	1/13	NA	NA
Kuwano et al. ²² 1997	Medical history, physical examination, laboratory tests, chest radiography, pulmonary function tests, and pulmonary biopsies	7	ADV	lung	PCR and ISH	3/19	7/30	52.61	24/25
Egan et al. ²³ 1995	Typical histological and clinical picture	8	EBV	lung	IHC	14/20	2/21	59.54	31/10
Ueda et al. ²⁴ 1992	Symptoms of progressive dyspnea and cough; audible fine crackles; chest radiograph; pulmonary function tests; no manifestations compatible with other interstitial lung diseases	8	HCV	serum	ELISA and RIBA	19/66	346/9464	61.5	46/20
Meliconi et al. ²⁵ 1996	Clinical, radiological, physiological, and histological grounds	8	HCV	serum	ELISA and RIBA	8/60	22/4744	62.34	106/84
Matsuyama et al. ²⁶ 2003	Chest radiograph, physical examination, lung function, pathological findings and no known diseases	7	HTLV-1	serum	serologic test and western blot	18/72	772/4782	62.4	NA

Vergnon et al. ²⁷	Clinical, chest X-ray, lung function, and histology	serologic test							
1984		8	EBV	serum	(IgM against VCA)	3/13	2/12	62.4	11/14
Barbera et al. ²⁸ 1992	Histopathology	8	EBV	lung	ISH	2/10	9/14	56.83	12/12
Kaan et al. ²⁹ 1997	Histopathology	8	EBV	lung	IHC	3/9	13/22	59.44	NA

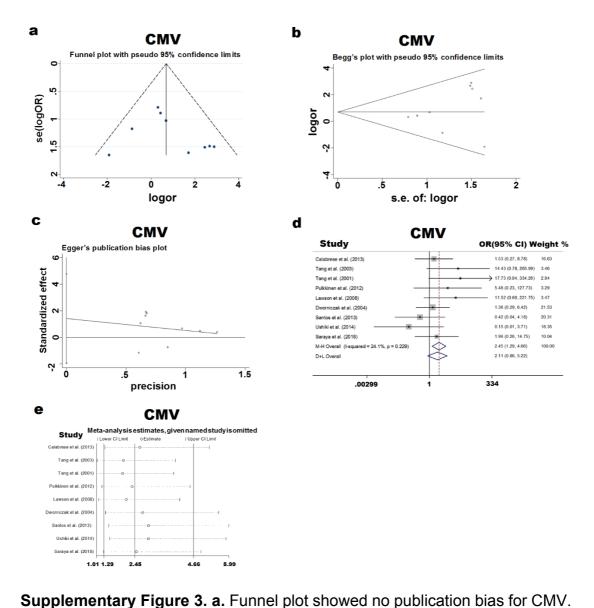
ATS, American Thoracic Society; ERS, European Respiratory Society; NOS, Newcastle-Ottawa Scale; y, year; M/F, male/female; NA, not available; IHC, immunohistochemistry; ISH, in situ hybridization; PCR, polymerase chain reaction; ELISA, enzyme linked immunosorbent assay; RIBA, recombinant immunoblotting assay; VCA, viral capsid antigen; EBV, Epstein-Barr virus; ADV, adenovirus; HCV, hepatitis C virus; HTLV, human T lymphotrophic virus



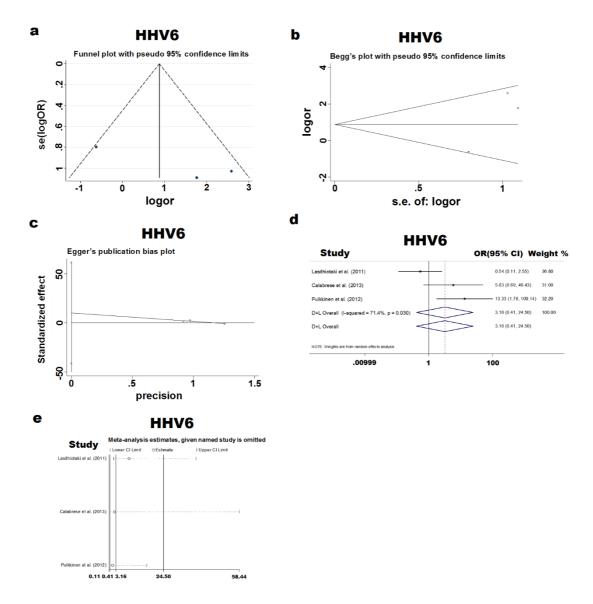
Supplementary Figure 1. a. Funnel plot showed a certain extent publication bias for all viruses. **b.** Begg's funnel plot showed a certain extent publication bias of OR for all viruses. **c.** Egger's funnel plot showed a certain extent publication bias of OR for all viruses. **d.** Begg's funnel plot using trim and fill method. Circle indicates included individual study and box the hypothetical study estimated by trim and fill method. **e.** The similarity between random and fixed effect models showed no significant influence on the pooled effect. **f.** Sensitive analysis of OR showed no significant difference.



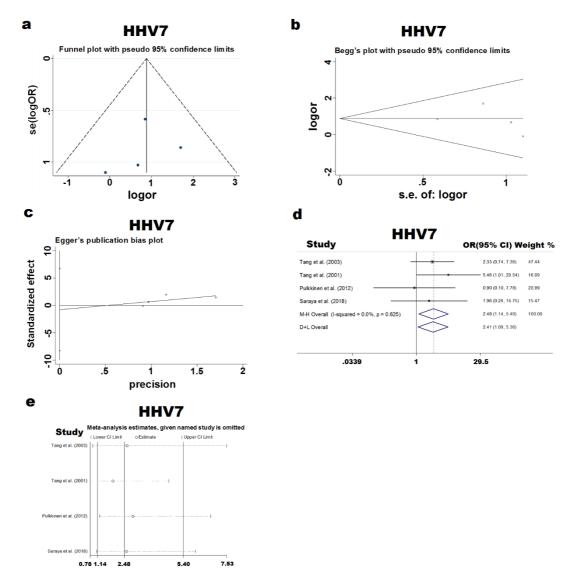
Supplementary Figure 2. a. Funnel plot showed a certain extent publication bias for EBV. **b.** Begg's funnel plot showed a certain extent publication bias of OR for EBV. **c.** Egger's funnel plot showed a certain extent publication bias of OR for EBV. **d.** Begg's funnel plot using trim and fill method. Circle indicates included individual study and box the hypothetical study estimated by trim and fill method. **e.** The similarity between random and fixed effect models showed no significant influence on the pooled effect. **f.** Sensitive analysis of OR showed no significant difference.



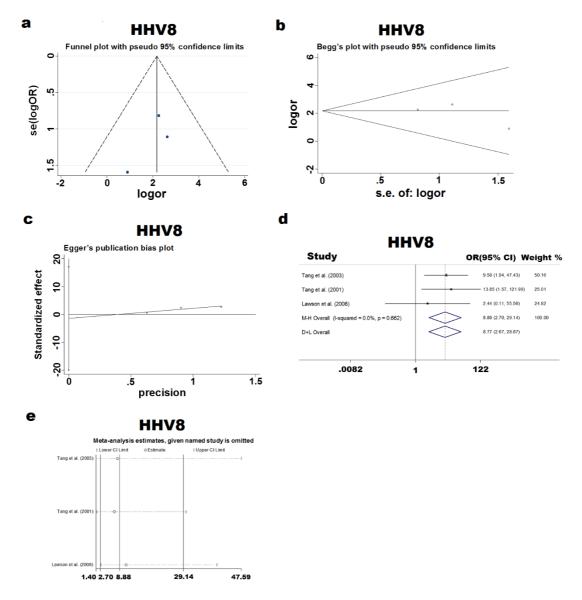
b. Begg's funnel plot showed no publication bias of OR for CMV. **c.** Egger's funnel plot showed no publication bias of OR for CMV. **d.** The similarity between random and fixed effect models showed no significant influence on the pooled effect. **e.** Sensitive analysis of OR showed no significant difference.



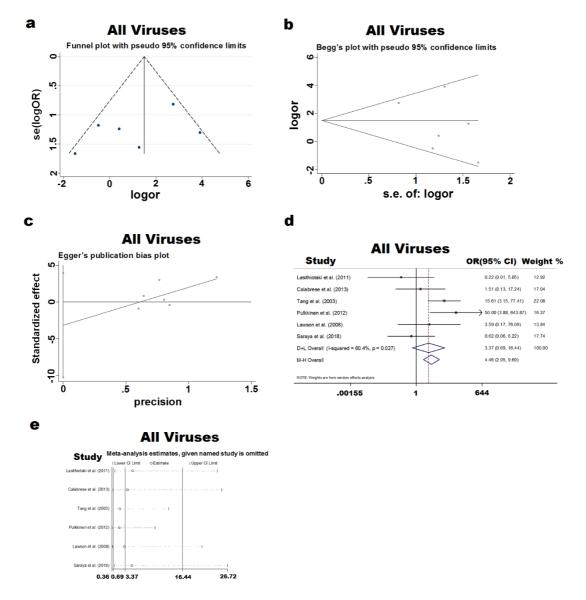
Supplementary Figure 4. a. Funnel plot showed no publication bias for HHV6. **b.** Begg's funnel plot showed no publication bias of OR for HHV6. **c.** Egger's funnel plot showed no publication bias of OR for HHV6. **d.** The similarity between random and fixed effect models showed no significant influence on the pooled effect. **e.** Sensitive analysis of OR showed no significant difference.



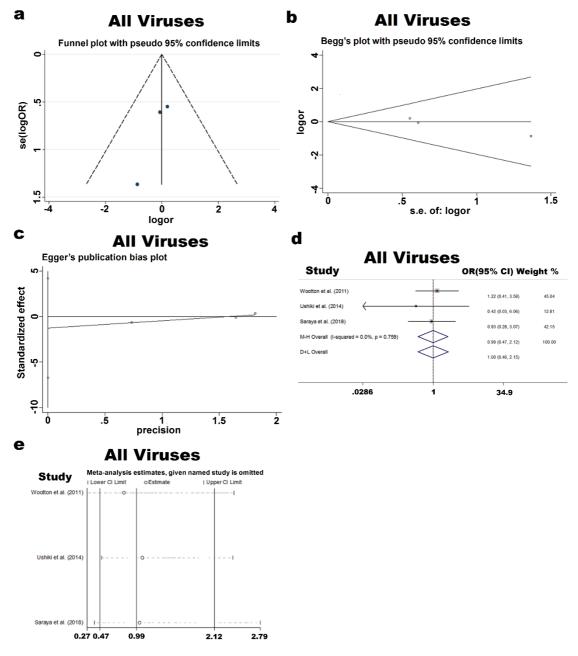
Supplementary Figure 5. a. Funnel plot showed no publication bias for HHV7. **b.** Begg's funnel plot showed no publication bias of OR for HHV7. **c.** Egger's funnel plot showed no publication bias of OR for HHV7. **d.** The similarity between random and fixed effect models showed no significant influence on the pooled effect. **e.** Sensitive analysis of OR showed no significant difference.



Supplementary Figure 6. a. Funnel plot showed no publication bias for HHV8. **b.** Begg's funnel plot showed no publication bias of OR for HHV8. **c.** Egger's funnel plot showed no publication bias of OR for HHV8. **d.** The similarity between random and fixed effect models showed no significant influence on the pooled effect. **e.** Sensitive analysis of OR showed no significant difference.



Supplementary Figure 7. a. Funnel plot showed no publication bias for multiple viral infections. **b.** Begg's funnel plot showed no publication bias of OR for multiple viral infections. **c.** Egger's funnel plot showed no publication bias of OR for multiple viral infections. **d.** The similarity between random and fixed effect models showed no significant influence on the pooled effect. **e.** Sensitive analysis of OR showed no significant difference.



Supplementary Figure 8. a. Funnel plot showed no publication bias in acute exacerbation of IPF for all viruses. **b.** Begg's funnel plot showed no publication bias of OR in acute exacerbation of IPF for all viruses. **c.** Egger's funnel plot showed no publication bias of OR in acute exacerbation of IPF for all viruses. **d.** The similarity between random and fixed effect models showed no significant influence on the pooled effect. **e.** Sensitive analysis of OR showed no significant difference.

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